



## GBA gene

glucosylceramidase beta

### Normal Function

The *GBA* gene provides instructions for making an enzyme called beta-glucocerebrosidase. This enzyme is active in lysosomes, which are structures inside cells that act as recycling centers. Lysosomes use digestive enzymes to break down toxic substances, digest bacteria that invade the cell, and recycle worn-out cell components. Based on these functions, enzymes in the lysosome are sometimes called housekeeping enzymes. Beta-glucocerebrosidase is a housekeeping enzyme that helps break down a large molecule called glucocerebroside into a sugar (glucose) and a simpler fat molecule (ceramide).

### Health Conditions Related to Genetic Changes

#### Gaucher disease

More than 200 mutations in the *GBA* gene have been identified in people with Gaucher disease, a disorder with varied features that affect many parts of the body. These mutations occur in both copies of the gene in each cell. Most of the *GBA* mutations responsible for Gaucher disease change a single protein building block (amino acid) in beta-glucocerebrosidase, altering the structure of the enzyme and preventing it from working normally. Other mutations delete or insert genetic material in the *GBA* gene or lead to the production of an abnormally short, nonfunctional version of the enzyme.

Mutations in the *GBA* gene greatly reduce or eliminate the activity of beta-glucocerebrosidase in cells. As a result, glucocerebroside is not broken down properly. This molecule and related substances can build up in white blood cells called macrophages in the spleen, liver, bone marrow, and other organs. The abnormal accumulation and storage of these substances damages tissues and organs, causing the characteristic features of Gaucher disease.

#### Parkinson disease

Changes in the *GBA* gene are also associated with Parkinson disease and parkinsonism, which are similar disorders that affect movement and balance. People with Gaucher disease have mutations in both copies of the *GBA* gene in each cell, while those with a mutation in just one copy of the gene are called carriers. People with Gaucher disease and people who are carriers of a *GBA* gene mutation have an increased risk of developing Parkinson disease or parkinsonism.

Symptoms of Parkinson disease and parkinsonism result from the loss of nerve cells that produce dopamine. Dopamine is a chemical messenger that transmits signals within the brain to produce smooth physical movements. It remains unclear how *GBA* gene mutations are related to these disorders. Studies suggest that changes in this gene may contribute to the faulty breakdown of toxic substances in nerve cells by impairing the function of lysosomes. Alternatively, the changes may increase the formation of abnormal protein deposits. As a result, toxic substances or protein deposits could accumulate and kill dopamine-producing nerve cells, leading to abnormal movements and balance problems.

### other disorders

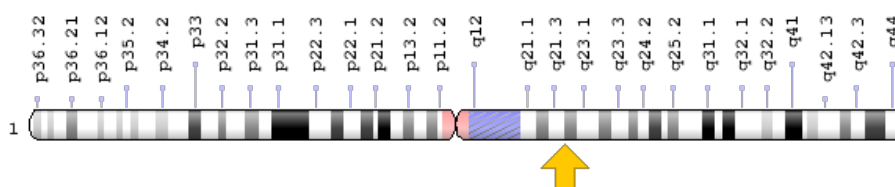
Research suggests an association between *GBA* gene mutations and a disorder called dementia with Lewy bodies. Lewy bodies are abnormal deposits of the protein alpha-synuclein that form in some dead or dying nerve cells. Specifically, they occur in nerve cells that produce the chemical messenger dopamine. The features of this disorder are variable, but symptoms typically include a loss of intellectual functions (dementia), visual hallucinations, and changes in alertness and attention. Affected individuals may have features characteristic of Parkinson disease such as trembling or rigidity of limbs, slow movement, and impaired balance and coordination. Lewy bodies are also a feature of Parkinson disease, but these abnormal deposits tend to be more widespread in the brain in dementia with Lewy bodies.

Carriers of *GBA* gene mutations appear to have an increased risk of developing dementia with Lewy bodies, although it remains unclear how changes in this gene increase the risk. Researchers speculate that mutations in the *GBA* gene alter the structure of beta-glucocerebrosidase and impair the function of lysosomes. As a result, alpha-synuclein may not be processed properly, allowing the formation of Lewy bodies.

### **Chromosomal Location**

Cytogenetic Location: 1q22, which is the long (q) arm of chromosome 1 at position 22

Molecular Location: base pairs 155,234,448 to 155,244,862 on chromosome 1 (Homo sapiens Annotation Release 108, GRCh38.p7) (NCBI)



Credit: Genome Decoration Page/NCBI

## Other Names for This Gene

- Acid beta-glucosidase
- Alglucerase
- beta-D-glucosyl-N-acylsphingosine glucohydrolase
- Beta-glucocerebrosidase
- D-Glucosyl-N-acylsphingosine glucosylhydrolase
- GBA1
- GLCM\_HUMAN
- GLUC
- Glucocerebrosidase
- Glucocerebroside beta-Glucosidase
- glucosidase, beta, acid
- glucosidase, beta; acid (includes glucosylceramidase)
- glucosphingosine glucosylhydrolase
- Glucosylceramidase
- Glucosylceramide beta-Glucosidase
- Imiglucerase

## Additional Information & Resources

### Educational Resources

- HuGENet Case Study: Glucocerebrosidase Gene Mutations and Parkinson's Disease  
<https://www.cdc.gov/genomics/hugenet/CaseStudy/PARKINSON/PARKview.htm>
- National Institute of Neurological Disorders and Stroke: Dementia with Lewy Bodies  
<https://www.ninds.nih.gov/Disorders/All-Disorders/Dementia-Lewy-Bodies-Information-Page>
- NIH News Release: Researchers Uncover Genetic Clues to a Common Form of Age-Related Dementia (July 17, 2006)  
<https://www.genome.gov/19517231/>

### GeneReviews

- Gaucher Disease  
<https://www.ncbi.nlm.nih.gov/books/NBK1269>

### Scientific Articles on PubMed

- PubMed  
<https://www.ncbi.nlm.nih.gov/pubmed?term=%28%28GBA%5BTI%5D%29+OR+%28beta+glucosidase%5BTIAB%5D%29+OR+%28glucosylceramidase%5BTIAB%5D%29+OR+%28Glucocerebrosidase%5BTIAB%5D%29%29+AND+%28%28Genes%5BMH%5D%29+OR+%28Genetic+Phenomena%5BMH%5D%29%29+AND+%28english%5Bla%5D+AND+human%5Bmh%5D+AND+%22last+720+days%22%5Bdp%5D>

### OMIM

- DEMENTIA, LEWY BODY  
<http://omim.org/entry/127750>
- GLUCOSIDASE, BETA, ACID  
<http://omim.org/entry/606463>

### Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology  
[http://atlasgeneticsoncology.org/Genes/GC\\_GBA.html](http://atlasgeneticsoncology.org/Genes/GC_GBA.html)
- ClinVar  
<https://www.ncbi.nlm.nih.gov/clinvar?term=GBA%5Bgene%5D>
- HGNC Gene Symbol Report  
[http://www.genenames.org/cgi-bin/gene\\_symbol\\_report?q=data/hgnc\\_data.php&hgnc\\_id=4177](http://www.genenames.org/cgi-bin/gene_symbol_report?q=data/hgnc_data.php&hgnc_id=4177)
- NCBI Gene  
<https://www.ncbi.nlm.nih.gov/gene/2629>
- PDGene  
<http://www.pdgene.org/view?gene=GBA>
- UniProt  
<http://www.uniprot.org/uniprot/P04062>

### **Sources for This Summary**

- Aharon-Peretz J, Rosenbaum H, Gershoni-Baruch R. Mutations in the glucocerebrosidase gene and Parkinson's disease in Ashkenazi Jews. N Engl J Med. 2004 Nov 4;351(19):1972-7.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/15525722>
- Charrow J, Andersson HC, Kaplan P, Kolodny EH, Mistry P, Pastores G, Prakash-Cheng A, Rosenbloom BE, Scott CR, Wappner RS, Weinreb NJ. Enzyme replacement therapy and monitoring for children with type 1 Gaucher disease: consensus recommendations. J Pediatr. 2004 Jan;144(1):112-20. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/14722528>

- Clark LN, Kartsaklis LA, Wolf Gilbert R, Dorado B, Ross BM, Kisselev S, Verbitsky M, Mejia-Santana H, Cote LJ, Andrews H, Vonsattel JP, Fahn S, Mayeux R, Honig LS, Marder K. Association of glucocerebrosidase mutations with dementia with lewy bodies. *Arch Neurol*. 2009 May;66(5):578-83. doi: 10.1001/archneurol.2009.54.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/19433657>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2758782/>
- Clark LN, Ross BM, Wang Y, Mejia-Santana H, Harris J, Louis ED, Cote LJ, Andrews H, Fahn S, Waters C, Ford B, Frucht S, Ottman R, Marder K. Mutations in the glucocerebrosidase gene are associated with early-onset Parkinson disease. *Neurology*. 2007 Sep 18;69(12):1270-7.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/17875915>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3624967/>
- Cox TM. Gaucher disease: understanding the molecular pathogenesis of sphingolipidoses. *J Inherit Metab Dis*. 2001;24 Suppl 2:106-21; discussion 87-8. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/11758671>
- Germain DP. Gaucher's disease: a paradigm for interventional genetics. *Clin Genet*. 2004 Feb;65(2):77-86. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/14984463>
- Goker-Alpan O, Giasson BI, Eblan MJ, Nguyen J, Hurtig HI, Lee VM, Trojanowski JQ, Sidransky E. Glucocerebrosidase mutations are an important risk factor for Lewy body disorders. *Neurology*. 2006 Sep 12;67(5):908-10. Epub 2006 Jun 21.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/16790605>
- Jmoudiak M, Futerman AH. Gaucher disease: pathological mechanisms and modern management. *Br J Haematol*. 2005 Apr;129(2):178-88. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/15813845>
- Mata IF, Samii A, Schneer SH, Roberts JW, Griffith A, Leis BC, Schellenberg GD, Sidransky E, Bird TD, Leverenz JB, Tsuang D, Zabetian CP. Glucocerebrosidase gene mutations: a risk factor for Lewy body disorders. *Arch Neurol*. 2008 Mar;65(3):379-82. doi: 10.1001/archneurol.2007.68.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/18332251>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2826203/>
- Orvisky E, Park JK, Parker A, Walker JM, Martin BM, Stubblefield BK, Uyama E, Tayebi N, Sidransky E. The identification of eight novel glucocerebrosidase (GBA) mutations in patients with Gaucher disease. *Hum Mutat*. 2002 Apr;19(4):458-9.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/11933202>
- Pelled D, Trajkovic-Bodennec S, Lloyd-Evans E, Sidransky E, Schiffmann R, Futerman AH. Enhanced calcium release in the acute neuronopathic form of Gaucher disease. *Neurobiol Dis*. 2005 Feb;18(1):83-8.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/15649698>

- Sidransky E, Nalls MA, Aasly JO, Aharon-Peretz J, Annesi G, Barbosa ER, Bar-Shira A, Berg D, Bras J, Brice A, Chen CM, Clark LN, Condroyer C, De Marco EV, Dürr A, Eblan MJ, Fahn S, Farrer MJ, Fung HC, Gan-Or Z, Gasser T, Gershoni-Baruch R, Giladi N, Griffith A, Gurevich T, Januario C, Kropp P, Lang AE, Lee-Chen GJ, Lesage S, Marder K, Mata IF, Mirelman A, Mitsui J, Mizuta I, Nicoletti G, Oliveira C, Ottman R, Orr-Urtreger A, Pereira LV, Quattrone A, Rogaeva E, Rolfs A, Rosenbaum H, Rozenberg R, Samii A, Samaddar T, Schulte C, Sharma M, Singleton A, Spitz M, Tan EK, Tayebi N, Toda T, Troiano AR, Tsuji S, Wittstock M, Wolfsberg TG, Wu YR, Zabetian CP, Zhao Y, Ziegler SG. Multicenter analysis of glucocerebrosidase mutations in Parkinson's disease. *N Engl J Med*. 2009 Oct 22;361(17):1651-61. doi: 10.1056/NEJMoa0901281.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/19846850>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC2856322/>
- Sidransky E. Gaucher disease: complexity in a "simple" disorder. *Mol Genet Metab*. 2004 Sep-Oct; 83(1-2):6-15. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/15464415>
- Sidransky E. Heterozygosity for a Mendelian disorder as a risk factor for complex disease. *Clin Genet*. 2006 Oct;70(4):275-82. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/16965318>
- Velayati A, Yu WH, Sidransky E. The role of glucocerebrosidase mutations in Parkinson disease and Lewy body disorders. *Curr Neurol Neurosci Rep*. 2010 May;10(3):190-8. doi: 10.1007/s11910-010-0102-x. Review.  
*Citation on PubMed:* <https://www.ncbi.nlm.nih.gov/pubmed/20425034>  
*Free article on PubMed Central:* <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3529411/>

Reprinted from Genetics Home Reference:  
<https://ghr.nlm.nih.gov/gene/GBA>

Reviewed: May 2012

Published: March 21, 2017

Lister Hill National Center for Biomedical Communications  
U.S. National Library of Medicine  
National Institutes of Health  
Department of Health & Human Services